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Display

Range: from  to  ☐ Reverse complemented strand Features: ☐ SNP ☒ CDD

☐ 1: [NM\\_002771](#). Reports Homo sapiens prot...[gi:21536451]

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LOCUS NM\_002771 807 bp mRNA linear PRI 27-NOV-2005  
 DEFINITION Homo sapiens protease, serine, 3 (mesotrypsin) (PRSS3), mRNA.  
 ACCESSION NM\_002771 NM\_007343  
 VERSION NM\_002771.2 GI:21536451  
 KEYWORDS .  
 SOURCE Homo sapiens (human)  
 ORGANISM Homo sapiens  
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;  
 Hominidae; Homo.

REFERENCE 1 (bases 1 to 807)  
 AUTHORS Marsit,C.J., Okpukpara,C., Danaee,H. and Kelsey,K.T.  
 TITLE Epigenetic silencing of the PRSS3 putative tumor suppressor gene in non-small cell lung cancer  
 JOURNAL Mol. Carcinog. 44 (2), 146-150 (2005)  
 PUBMED [16013053](#)  
 REMARK GeneRIF: we determined the promoter hypermethylation status of PRSS3 in a case series study of primary NSCLC, and found methylation of this gene to be common, occurring in 53% (86 of 166) of tumors examined.

REFERENCE 2 (bases 1 to 807)  
 AUTHORS Nemoda,Z., Teich,N., Hugenberg,C. and Sahin-Toth,M.  
 TITLE Genetic and biochemical characterization of the E32del polymorphism in human mesotrypsinogen  
 JOURNAL FEBS Lett. 5 (2-3), 273-278 (2005)  
 PUBMED [15855826](#)  
 REMARK GeneRIF: The results classify E32del mesotrypsinogen as a frequent polymorphic variant, which is not associated with chronic alcoholic pancreatitis

REFERENCE 3 (bases 1 to 807)  
 AUTHORS Szmola,R., Kukor,Z. and Sahin-Toth,M.  
 TITLE Human mesotrypsin is a unique digestive protease specialized for the degradation of trypsin inhibitors  
 JOURNAL J. Biol. Chem. 278 (49), 48580-48589 (2003)  
 PUBMED [14507909](#)  
 REMARK GeneRIF: biological function of human mesotrypsin is digestive degradation of trypsin inhibitors

REFERENCE 4 (bases 1 to 807)  
 AUTHORS Katona,G., Berglund,G.I., Hajdu,J., Graf,L. and Szilagyi,L.  
 TITLE Crystal structure reveals basis for the inhibitor resistance of human brain trypsin  
 JOURNAL J. Mol. Biol. 315 (5), 1209-1218 (2002)  
 PUBMED [11827488](#)

REMARK GeneRIF: X-ray structure in complex with the inhibitor benzamidine at 1.7 A resolution; crystal structure reveals basis for inhibitor resistance

REFERENCE 5 (bases 1 to 807)  
AUTHORS Nyaruhucha,C.N., Kito,M. and Fukuoka,S.I.  
TITLE Identification and expression of the cDNA-encoding human mesotrypsin(ogen), an isoform of trypsin with inhibitor resistance  
JOURNAL J. Biol. Chem. 272 (16), 10573-10578 (1997)  
PUBMED [9099703](#)

REFERENCE 6 (bases 1 to 807)  
AUTHORS Stubbs,M.T., Huber,R. and Bode,W.  
TITLE Crystal structures of factor Xa specific inhibitors in complex with trypsin: structural grounds for inhibition of factor Xa and selectivity against thrombin  
JOURNAL FEBS Lett. 375 (1-2), 103-107 (1995)  
PUBMED [7498454](#)

REFERENCE 7 (bases 1 to 807)  
AUTHORS Wiegand,U., Corbach,S., Minn,A., Kang,J. and Muller-Hill,B.  
TITLE Cloning of the cDNA encoding human brain trypsinogen and characterization of its product  
JOURNAL Gene 136 (1-2), 167-175 (1993)  
PUBMED [8294000](#)

REFERENCE 8 (bases 1 to 807)  
AUTHORS Tani,T., Kawashima,I., Mita,K. and Takiguchi,Y.  
TITLE Nucleotide sequence of the human pancreatic trypsinogen III cDNA  
JOURNAL Nucleic Acids Res. 18 (6), 1631 (1990)  
PUBMED [2326201](#)

REFERENCE 9 (bases 1 to 807)  
AUTHORS Rinderknecht,H., Renner,I.G., Abramson,S.B. and Carmack,C.  
TITLE Mesotrypsin: a new inhibitor-resistant protease from a zymogen in human pancreatic tissue and fluid  
JOURNAL Gastroenterology 86 (4), 681-692 (1984)  
PUBMED [6698368](#)

COMMENT REVIEWED REFSEQ: This record has been curated by NCBI staff. The reference sequence was derived from [X15505.1](#).  
On or before Dec 15, 2003 this sequence version replaced [gi:20428774](#), [gi:4506148](#).

Summary: This gene encodes a trypsinogen, which is a member of the trypsin family of serine proteases. This enzyme is expressed in the brain and pancreas and is resistant to common trypsin inhibitors. It is active on peptide linkages involving the carboxyl group of lysine or arginine. This gene is localized to the locus of T cell receptor beta variable orphans on chromosome 9. Additional transcript variants for this gene have been described, but their full length sequences have not been determined.  
COMPLETENESS: complete on the 3' end.

FEATURES

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ORIGIN

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